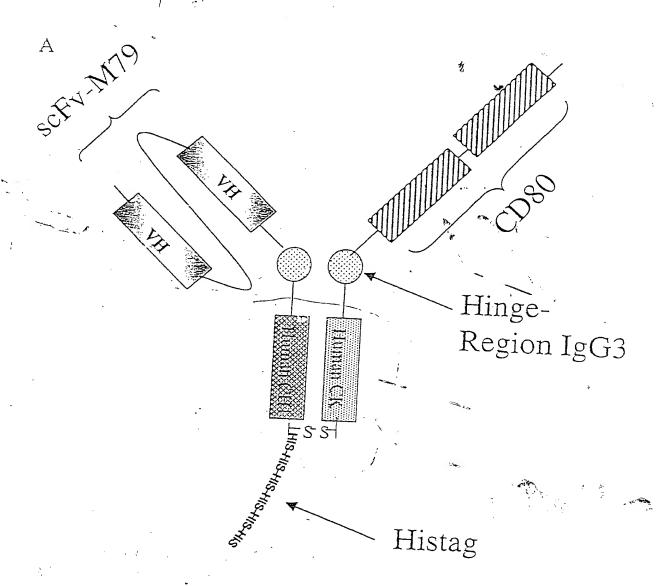
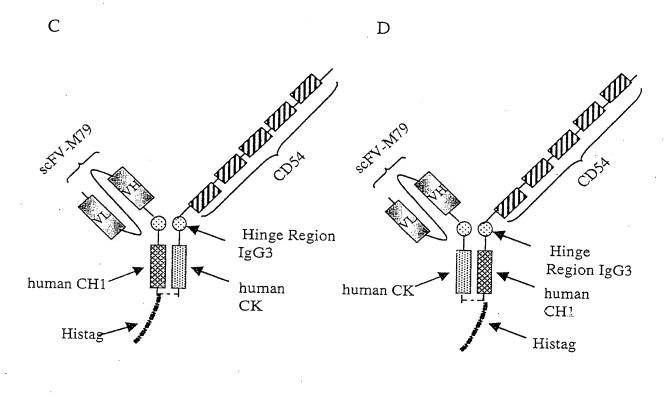
Figure 1

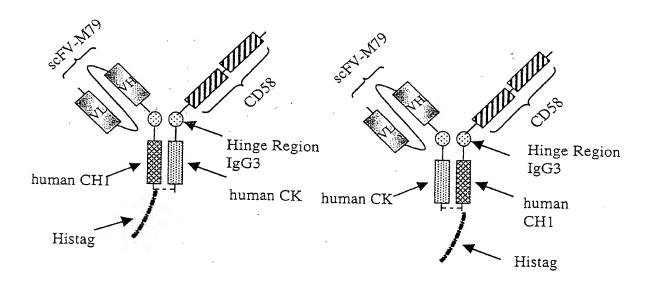


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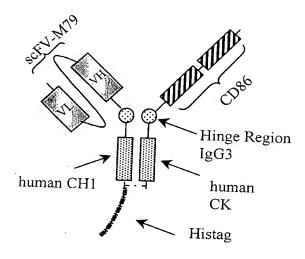


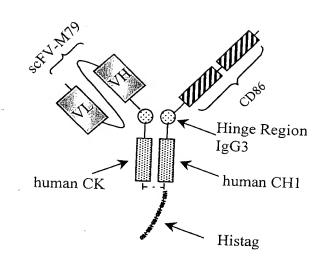
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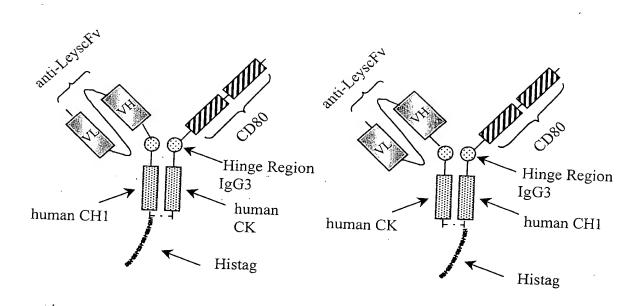


Figure 2

EcoRI

AsuII XbaIBspE1

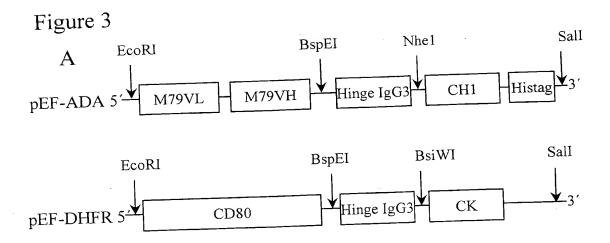
BamHI EcoRV XmaI

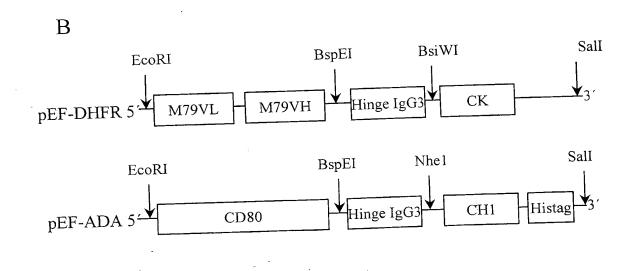
5 CTA GAA TTC TTC GAA TCC GGA GGT GGT GGA TCC GAT ATC CCC GGG

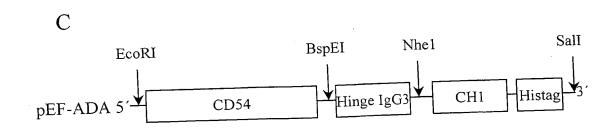
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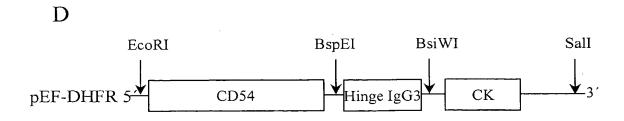
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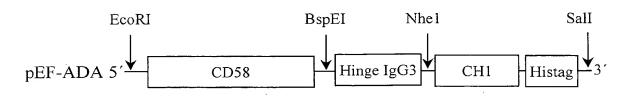




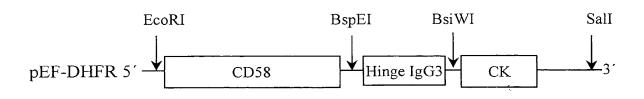




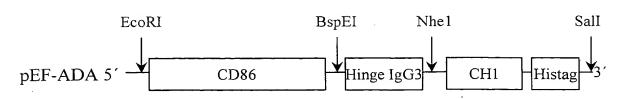
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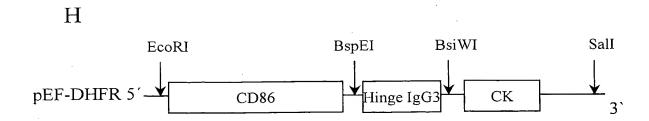


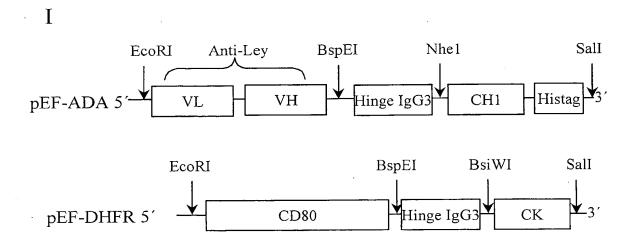
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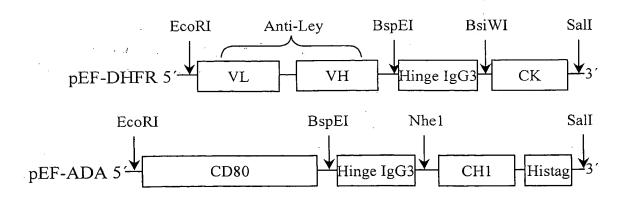






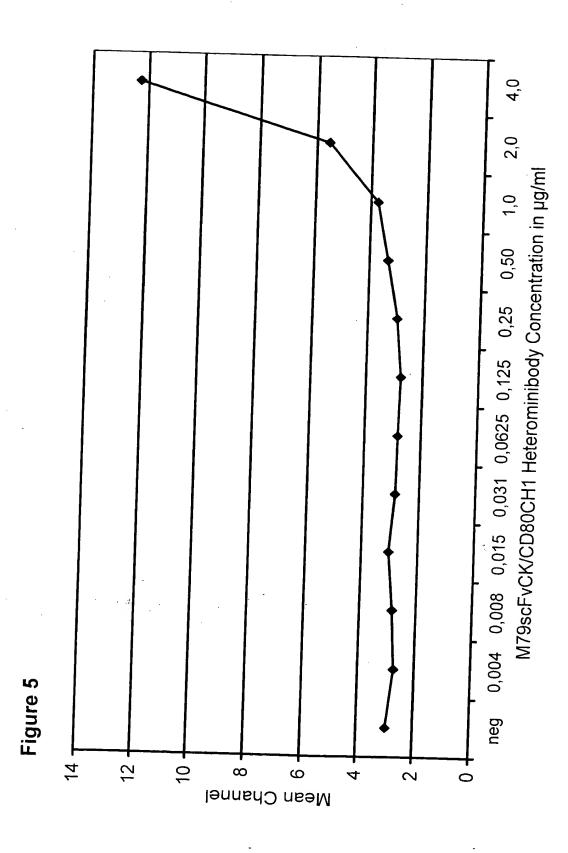
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Figure 6

		RI				19			28									5 5
5 '	GAAT	TCC	ACC	ATG	GG.A	TGG	AGC	TGT	ATC.	ATC	CTC	TTC	TTG	GTA	GCA	ACA	GCT	
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			64			73			82			91			100	•		109
	GGT	GTA			Car	_	CTT	GTG		CAG	G23		GC 2	CTC		AC 2	エこユ	
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	G	Ε	T	V	T	L	T	C	R	S	S	T	G	A	V	T	T	S
			172			181			1.90			199			208			217
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	N	¥	A	И	W	V	Q	Ξ	ĸ	Ď	D	H	L	F	T	ئ	L	I
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			226			235			244			253			252			271
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			280			289			298			307			316			325
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	A.A.A	CTC	GAG	GTC	CTA	GGT	GGT	GGT	GGT	TCT	GGC	GGC	GGC	GGC	TCC	GGT	GGT	GGT
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		496			505			514			523						541
AGC	CTG	TCC	ATC	ACA	TGC	ACC	ATC	TCA	GGG	TTC	TCA	TTA	ACT	AAG	TAT	GGT	G-17
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CAC	TGG	GTT	CGC	CAG	CCT	CCA	GG 3	220	GGT	CTG	GaG.	TGG	СТС	GTG	GTC	מידי ה	222
											C.A.G	-	CIG	010	010	AIA	ال (باري
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L	D	G	G	Τ.	S	ž	N	S	À	Ĺ	К	S	R	L	S	I	S
		658			667			676			685			694			70,3
AAG	GAC	658 AAC	TCC	AAG	667 AGC	CAA	GTT	676 TTC	TTA	AAA	685 ATG	AAC	AGT	694 CTC	CAA	ACT	703 GAT
AAG	GAC	658 AAC	TCC	AAG	667 AGC	CAA	GTT	676 TTC	TTA	A <u>A</u> A 	685 ATG	AAC 	AGT	694 CTC	CAA	A.CT	70,3 GAT
AAG K	GAC	658 AAC N	TCC	AAG	AGC	CAA Q	GTT V	676 TTC F	TTA	AAA K	685 ATG M	AAC N	AGT S	694 CTC L	CAA Q	ACT T	703 GAT
		AAC	TCC	AAG	AGC	C74	GTT	TTC	TTA	A.A.A.	ATG	AAC 	AGT	CTC	CAA	ACT	GAT
		AAC	TCC S	AAG K	AGC S	CAA Q	GTT V	TTC F	TTA L	A.A.A. 	ATG M	AAC N	AGT S	CTC L	CAA Q	A.CT T	GAT D
K	D	AAC N 712	TCC s	AAG K	AGC S 721	CAA Q	GTT V	TTC F 730	TTA L	A.A.A X	ATG M 73.9	AAC N	AGT S	CTC L 748	CAA Q	ACT T	GAT D 757
K	D	AAC N 712	TCC S	AAG K	AGC S 721	CAA Q	GTT V	TTC F 730	TTA L	A.A.A X	ATG M 73.9	AAC N	AGT S	CTC L 748	CAA Q	ACT T	GAT D 757
K	D	AAC N 712 GCC	TCC S ATG	AAG K	AGC S 721	CAA Q TGT	GTT V	TTC F 730	TTA L CAG	AAA K GAT	ATG M 739 AGA	AAC N TAC	AGT S S GAC	CTC L 748 GGT	CAA Q Q GGA	ATT	GAT O 757 GCT
K GAC	D ACA	AAC N 712 GCC	TCC S S ATG	AAG K TAC	AGC S 721 TAC	CAA Q TGT	GTT V GCC	TTC F 730 AGA	TTA L CAG	AAA K GAT	ATG M 73.9	AAC N	AGT S	CTC L 748	CAA Q	ACT T	GAT D 757
K GAC	D ACA	AAC N 712 GCC A	TCC S ATG M	AAG K TAC	AGC S 721 TAC Y	CAA Q TGT	GTT V GCC	TTC F 730 AGA R	TTA L CAG Q	K GAT	ATG M 73.9 AGA	AAC N TAC	AGT S S GAC	CTC L 748 GGT	CAA Q Q GGA	ATT	GAT O 757 GCT
K GAC	D ACA T	712 GCC A	TCC S ATG M	AAG K TAC Y	AGC S 721 TAC Y 775	CAA Q TGT	GTT V GCC A	730 AGA R	TTA L CAG	AAA K GAT	ATG M 73° AGA R	AAC N TAC Y	AGT S GAC D	CTC L 748 GGT	CAA Q Q GGA	ATT	GAT O 757 GCT
K GAC	D ACA T	712 GCC A	TCC S ATG M	AAG K TAC Y	AGC S 721 TAC Y 775	CAA Q TGT	GTT V GCC A	730 AGA R	TTA L CAG	AAA K GAT	ATG M 73° AGA R	AAC N TAC Y	AGT S GAC D	CTC L 748 GGT	CAA Q Q GGA	ATT	GAT O 757 GCT
K GAC	D ACA T	712 GCC A	TCC S ATG M	AAG K TAC Y	AGC S 721 TAC Y 775	CAA Q TGT	GTT V GCC A	730 AGA R	TTA L CAG	AAA K GAT	ATG M 73° AGA R	AAC N TAC Y	AGT S GAC D	CTC L 748 GGT	CAA Q Q GGA	ATT	GAT O 757 GCT

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Figure 7

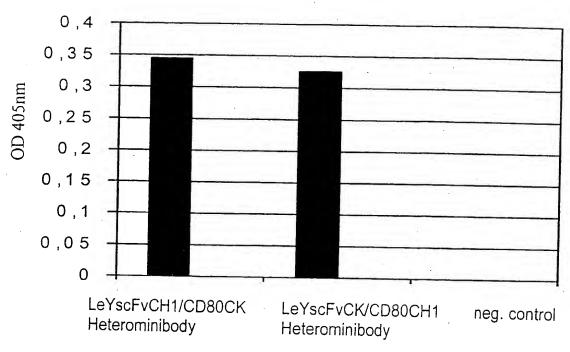
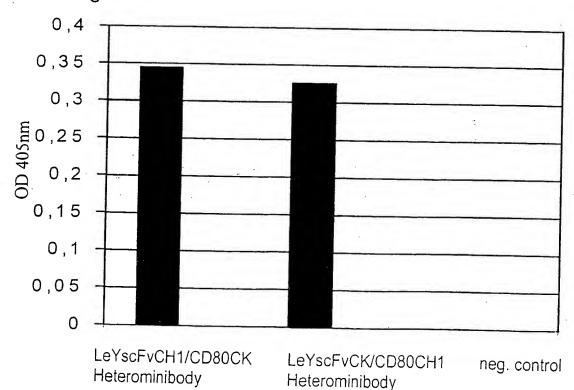
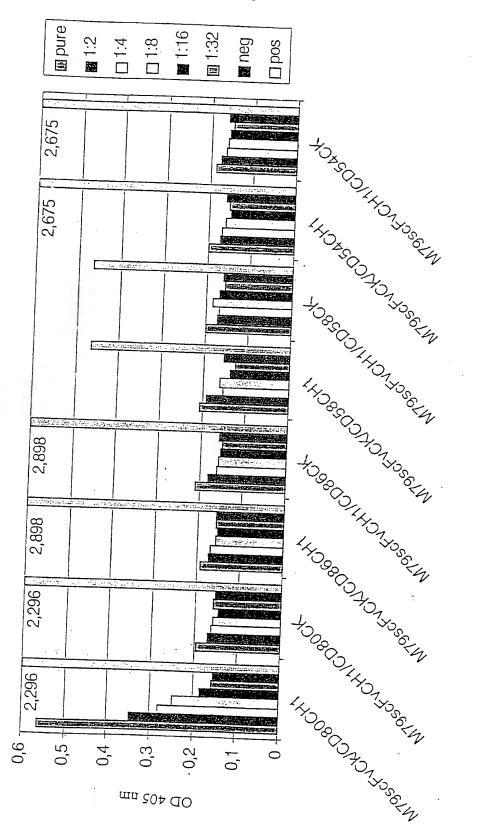


Figure 8



· YEAR SOLL



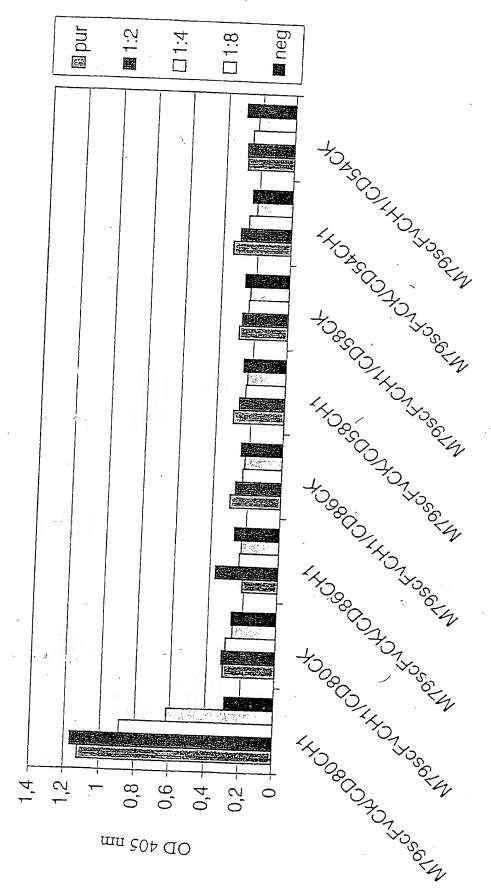
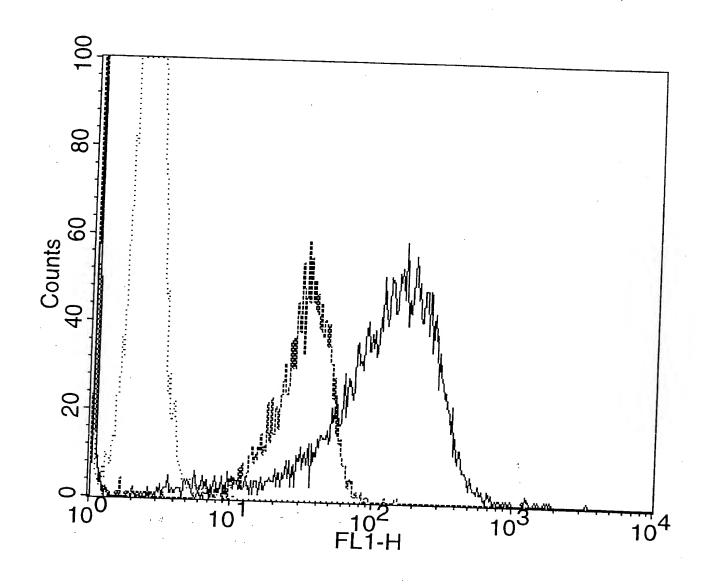


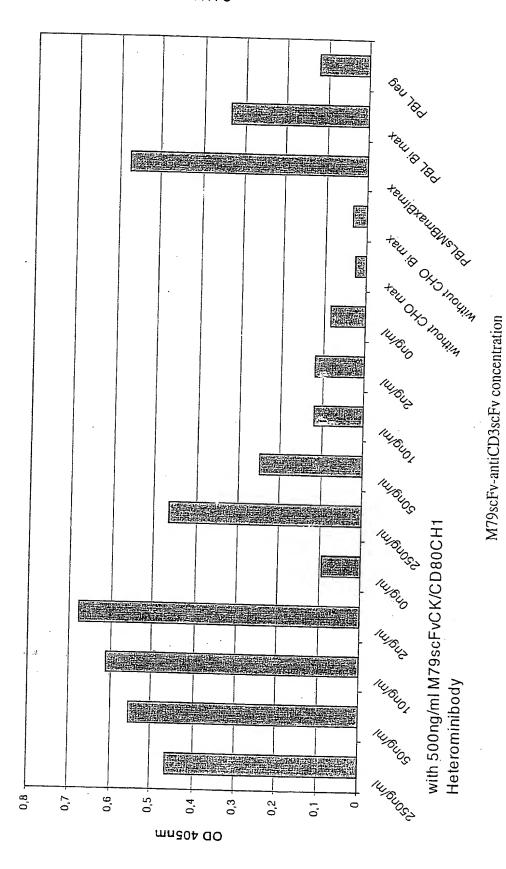
Figure 10

Figure 11

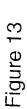


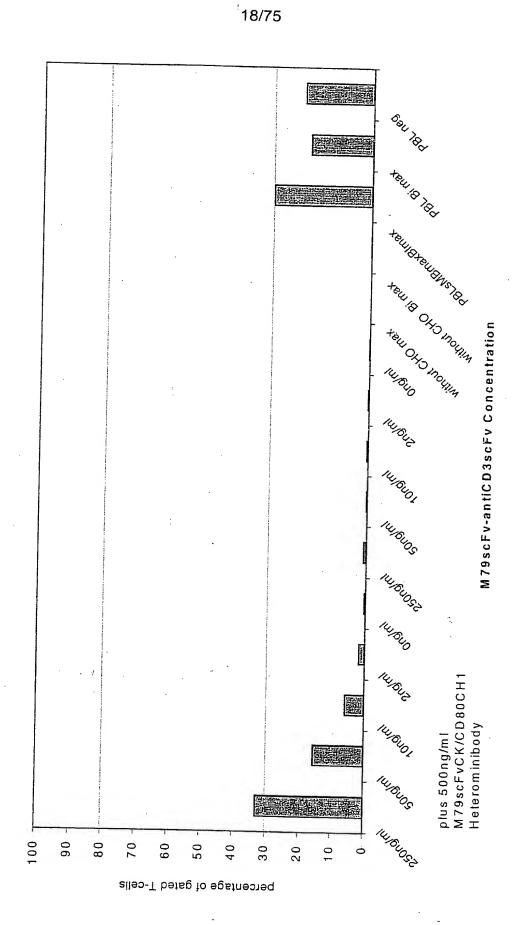
M79 on 17-1A transfected CHO cells
M79 on untransfected CHO cells
M79 on KATO cells

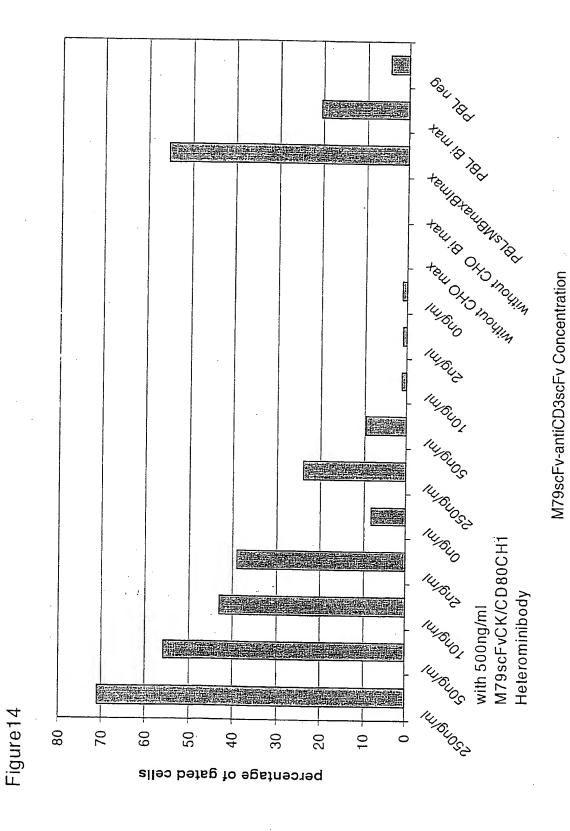


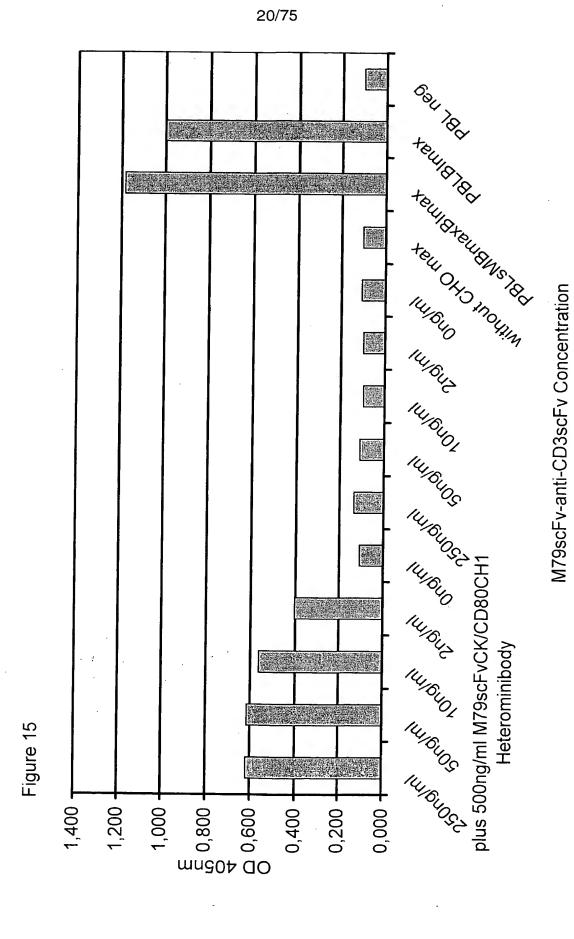


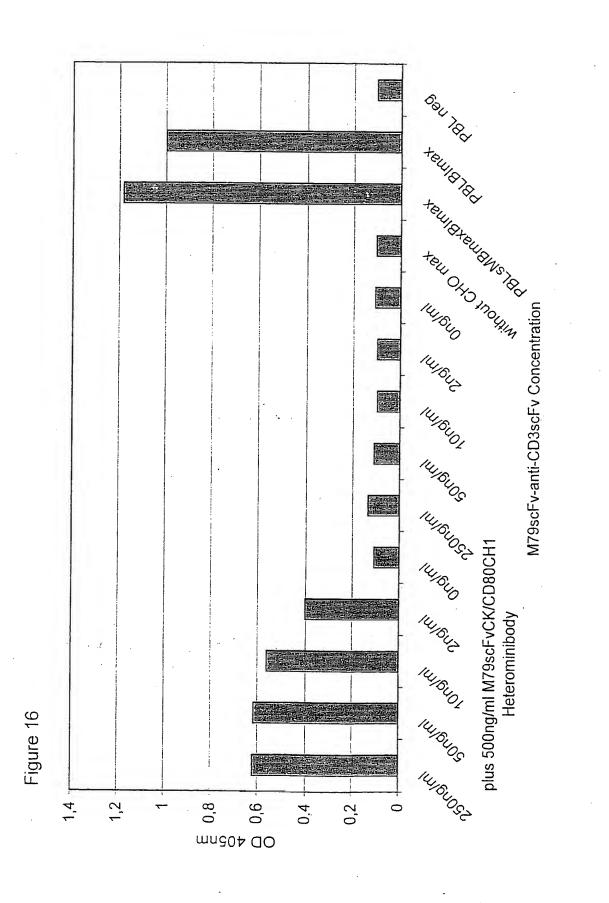
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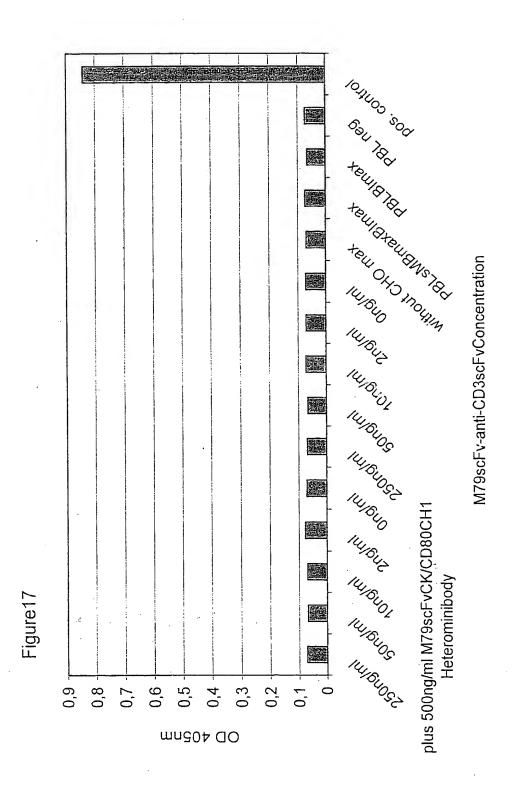


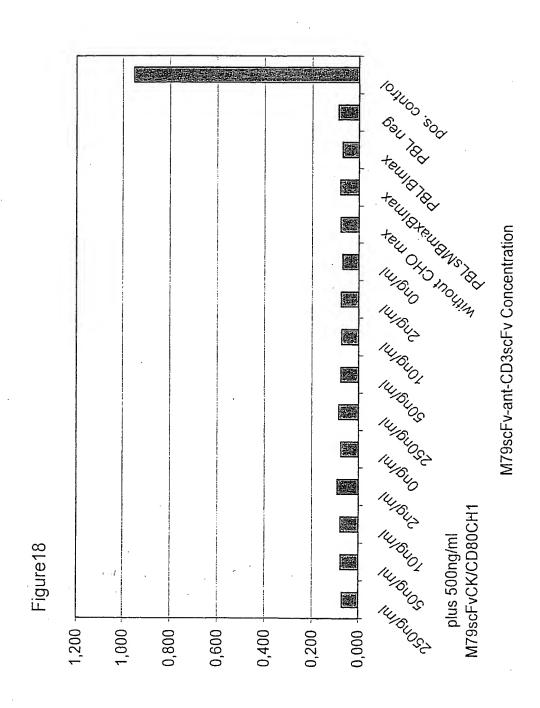




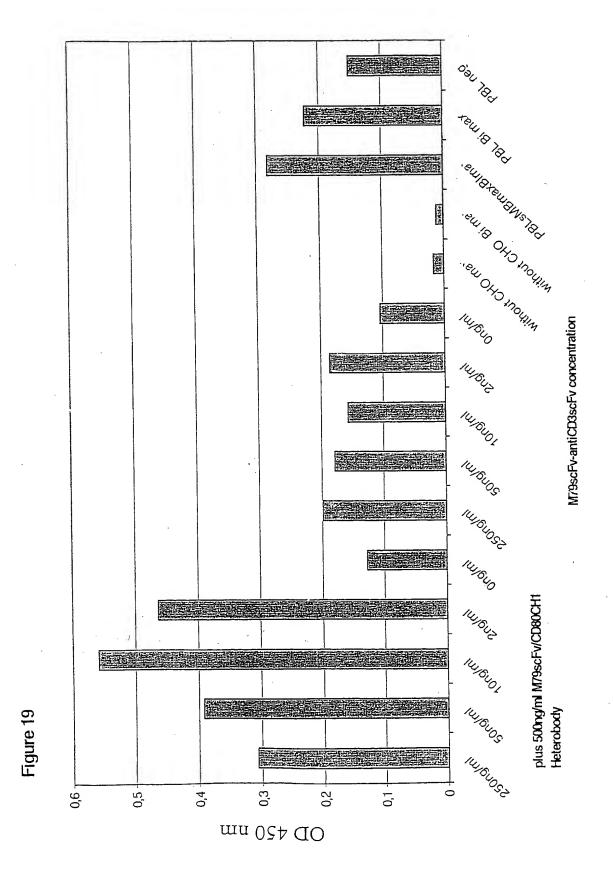


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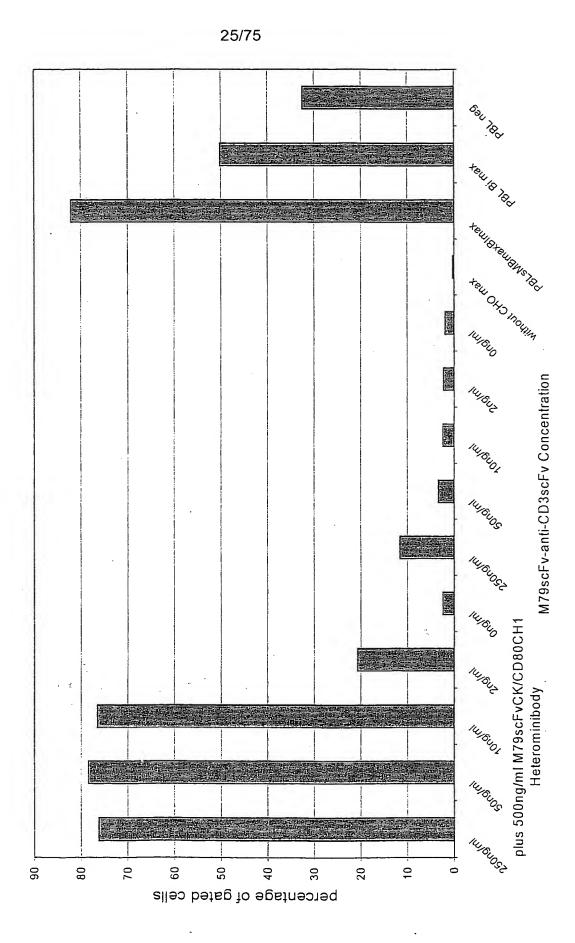


OD402nm

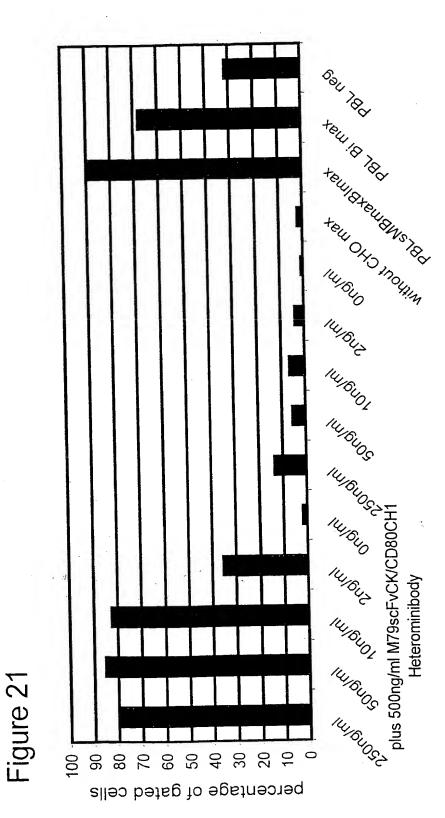


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M79scFv-anti-CD3scFv Concentration

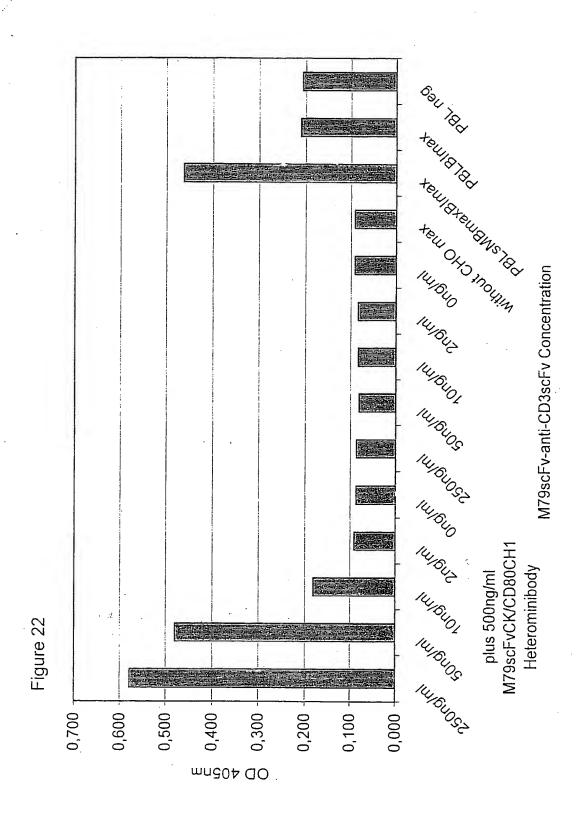
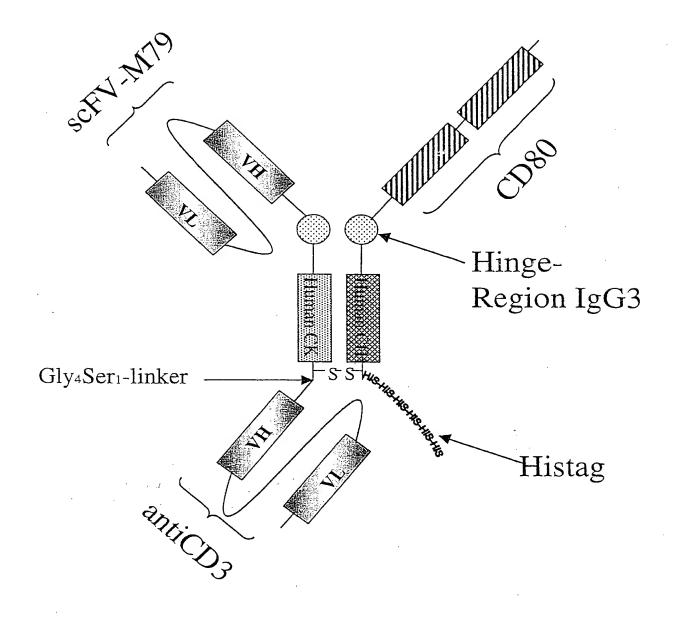


Figure 23



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Figure 24

Recombinant bifunctional single-chain protein

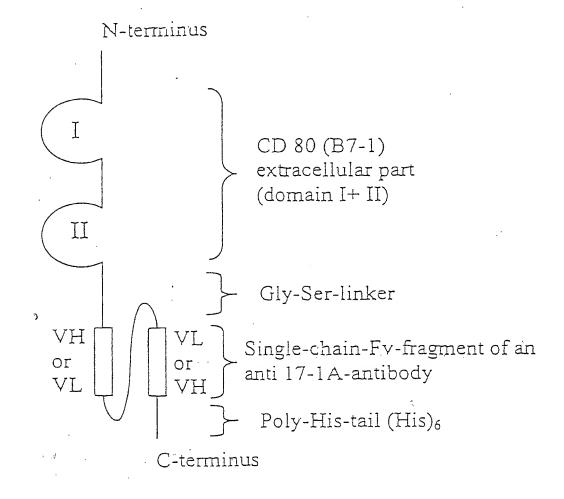
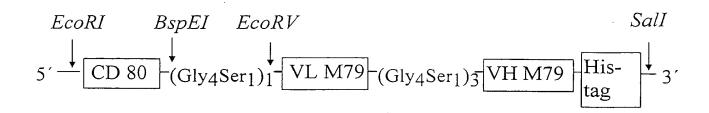
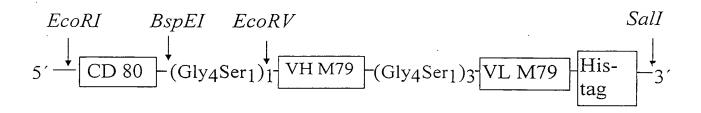
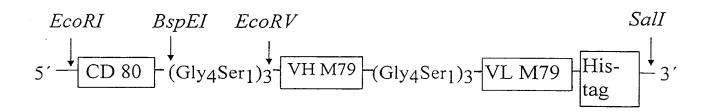


Figure 25

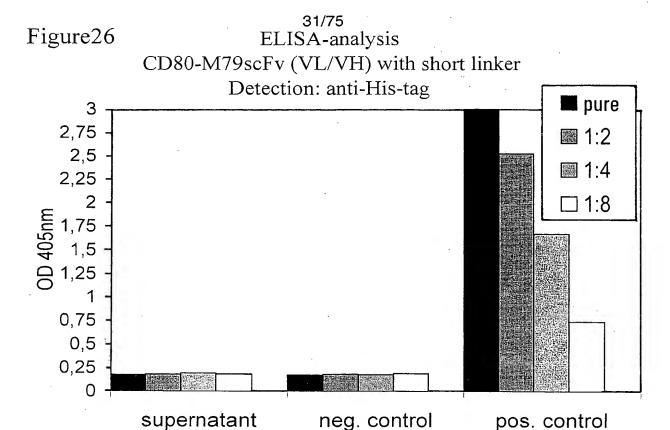






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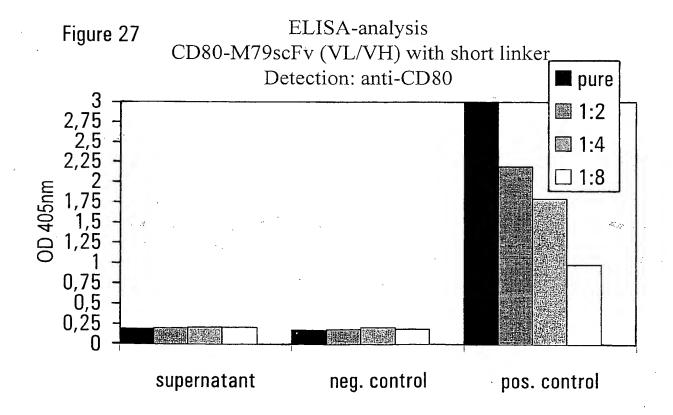
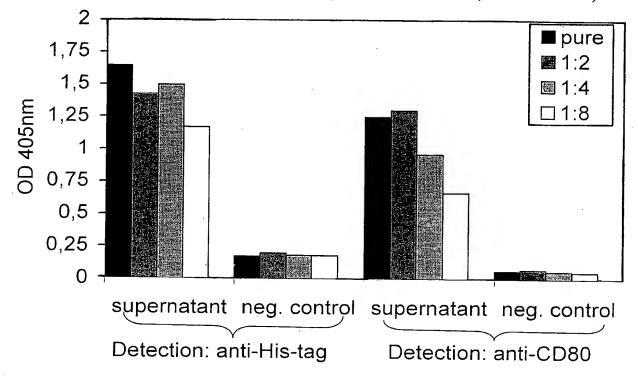


Figure 28

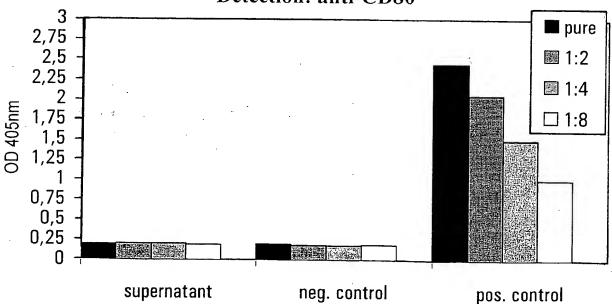
AND THE RESERVE

ELISA-analysis CD80-M79scFv (VL/VH) with short linker Detection: anti-His-tag or anti-CD80 (as indicated)



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Figure 29 ELISA-analysis
CD80-M79 scFv (VH/VL) with short linker
Detection: anti-CD80



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Figure 30 DNA-sequence of the double-stranded oligonucleotide designated **ACCGS15 BAM**

BspE1 T CCA CCA CCA AGG CCC CCA CCT CCA AGT CCG CCA CCA CCTAG 5

BamHI

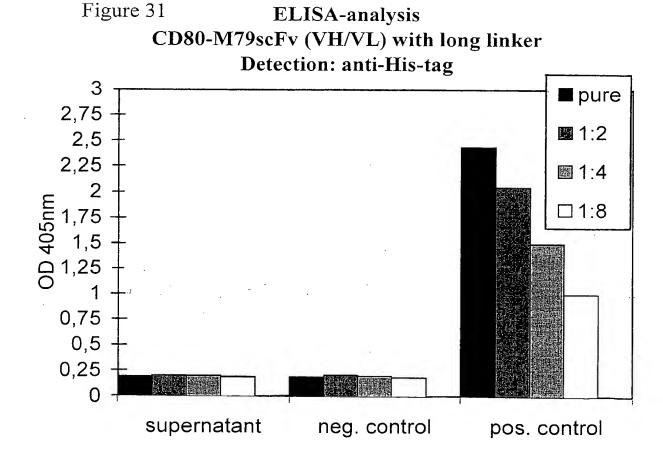
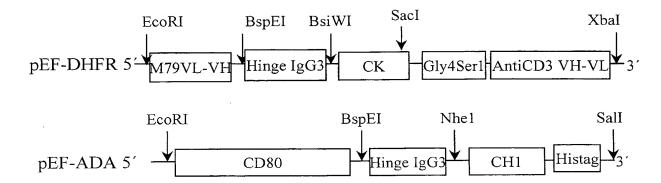


Figure 32



35/75 Figure 33 Phenotyp-Switch CD45 RA+/R0- naive T-cells Priming CD45 <u>RA-/R0+</u> effector/memory T-cells **PBMC** naive CD4+ T-cells separation day 0 $10^2 10^3 10^4$ CD45RO CD45RO day 6 5 R R CD45RO CD45RO

only primary signal: M79scFv-antiCD3scFv

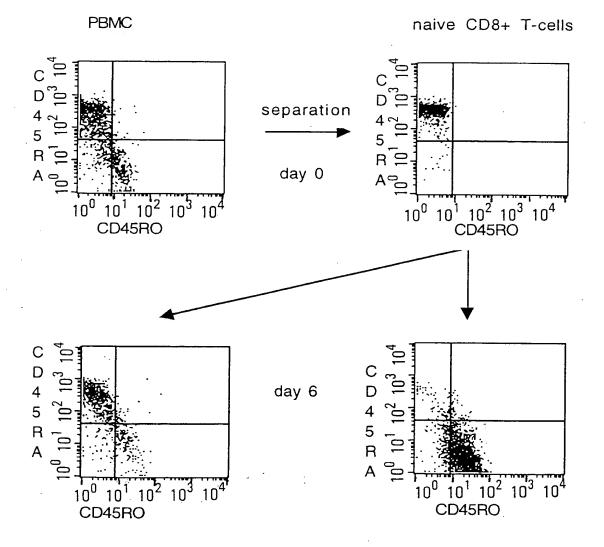
primary + costimulatory signal: M79scFv-antiCD3scFv and M79scFvCK/CD80CH1 Heterominibody

Figure 34 Phenotyp-Switch

CD45 <u>RA+/R0-</u> naive T-cells Priming

CD45 **RA-/R0+**

effector/memory T-cells



only primary signal: M79scFv-antiCD3scFv

primary + costimulatory signal: M79scFv-antiCD3scFv and M79scFvCK/CD80CH1 Heterominibody

(E.S.)

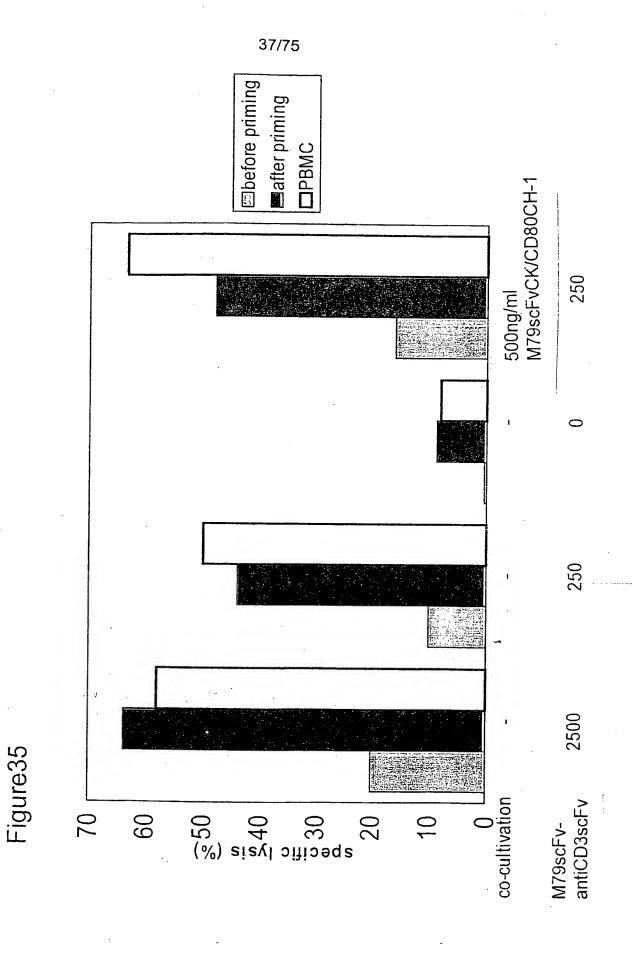


Figure 36

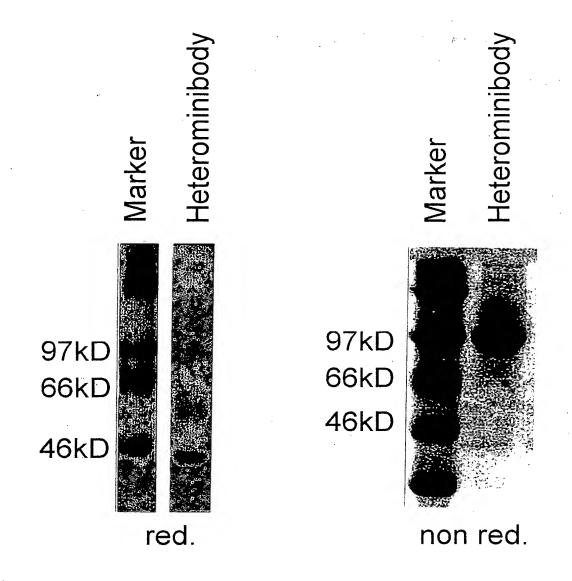
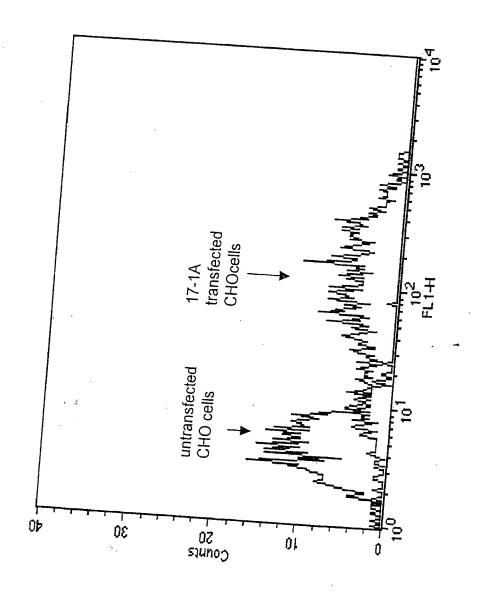
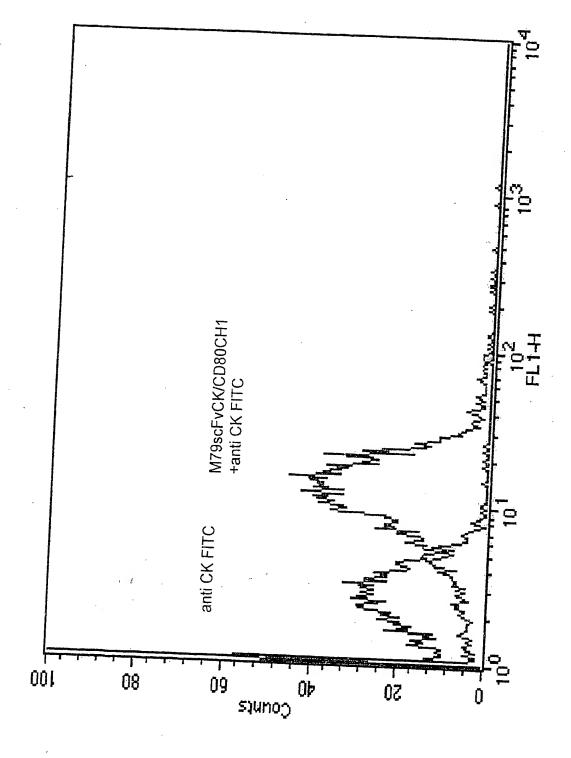




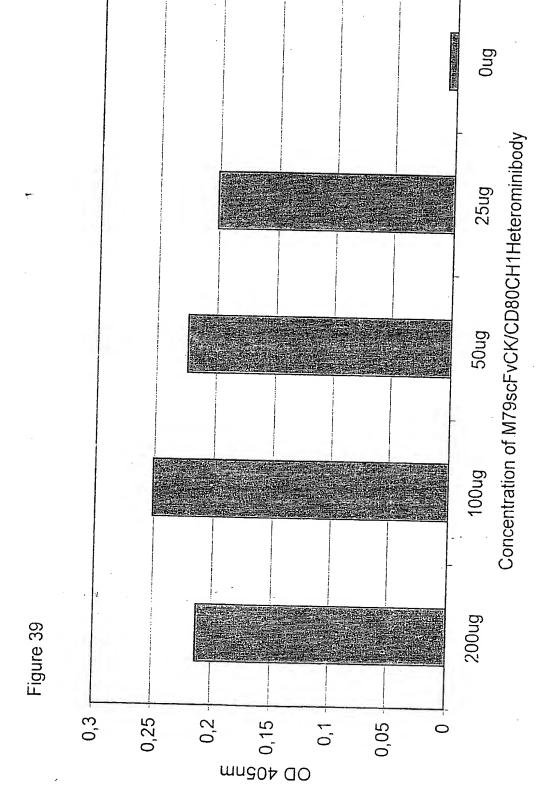
Figure 37

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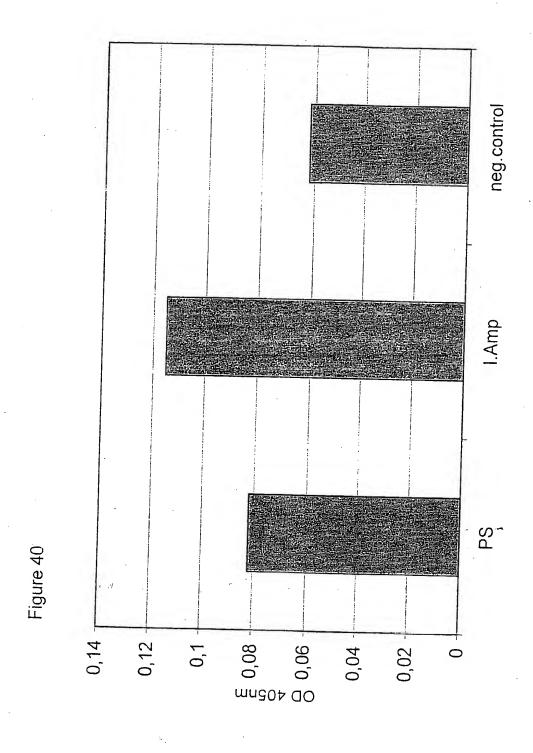




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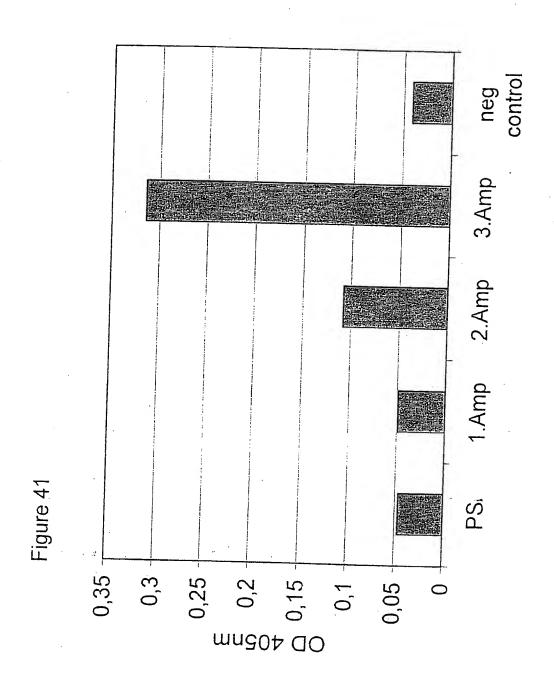
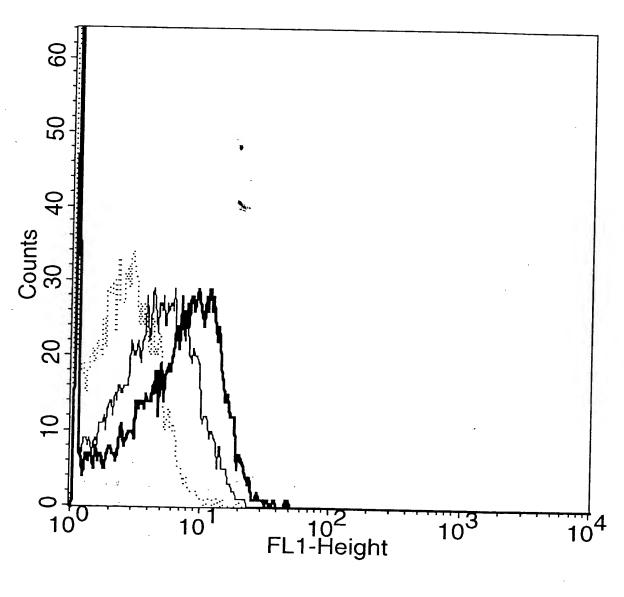


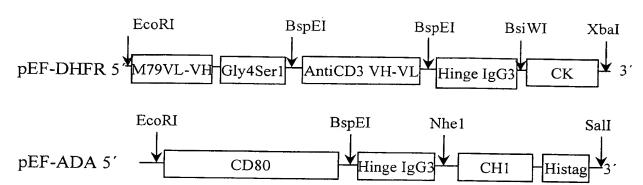
Figure 42



neg. control 50μg/ml Heterominibody 400µg/ml Heterominibody

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Fig 44



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Figure 45

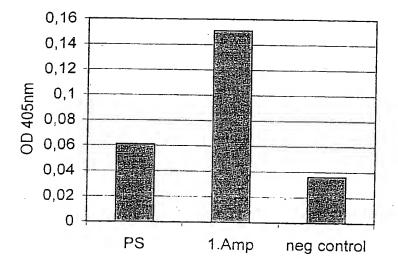


Figure 46

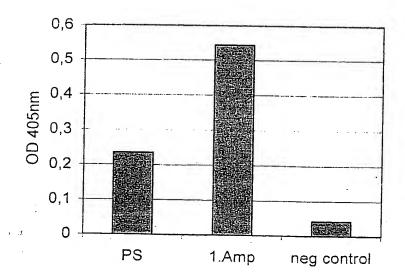


Figure 47:

	Figu	re 4,	<i>'</i> :															
5 '	کسے		9 ACC	ATG	GGA	18 TGG		TGT						GTA			GCT	J -
_																		
			•	M	G	W	S	С	Ŧ	. I	L	F	L	V	A	T·	A	T
	GGT	GTA	63 CAC	TCC	GAT	72 ATC	CAG	CTG	81 ACC	CAG	TCT	90 CAA	AAA	TTC	99 ATG	TCC	ACA	108 TCA
	G	V	. Н	S	D	I	Q	L	T	Q	S	Õ	K	F	M	s	\mathbf{T}	S
	GTÁ	GGA	117 GAC	AGG	GIC	126 AGC	GTC	ACC	135 TGC	AAG	GCC	144 AGT	CAG	AAT	153 GTG	GGT	ACT	162 AAT
	 V	G	D	R	V	s	V	T	C	K	A	s	Q	N	V	G	T	N
	GTA	GCC	171 TGG	TAT	CAA	180 CAG	AAA	CCA	189 GGG	CAA		198 CCT		GCA	207 CTG		TAC	216 TCG
	 V	 A		 Y	Q	Q	 К	P	G	Q	S	Ď	K	A	L	I	Y	s
	GCA		225 TAC		TAC	234	GGA	GTC	243 CCT	GAT	CGC	252 TTC	ACA	GGC	261 AGT	GGA	TCT	270 GGG
	 A	s	Y	R	Y	S	G	V	P	D	R	F	${f T}$	G	ŗs	G	S	G
	ACA	GAT	279 TTC	ACT	CTC	288 ACC	ATC	AGC	297 AAT	GTG	CAG	306 TCT	GAA	GAC	315 TIG	GCA	GAG	324 TAT
	 Т	 D	 F	 Т	- - -	 T	 I	 S	 N		Q		 E	 D	 L	 A	 E	 Y
	_	_	333	_		342			351			360			369			378
-	TTC	TGT	CAG	CAA	TAT	AAC	AGC	TAT	CCG	CTC	ACG	TTC	GGT	GCT	GGG	ACC	AAG	CIC
	F	С	Q	Q	Y	N	S	Y	P	L	${f T}$	F	G	A	G	T	K	L
	GAG	ATC	387 AAA	GGT	GGT	396 GGT		TCT			GGC	414 GGC	TCC	GGT	423 GGT	GGT	GGT	432 TCT
	E	I	K	. G	G	G	G	S	G	G	G	G	S	G	G	G	G	S
	CAG	GTG	441 AAA		CAG	450 GAG		GGA	459 CCT		CTA	468 GTG		CCC	477 TCA		AGC	486 CTG
	Q	V	K	L	· Q	E.	S	G	P	G	L	V	Q	P	s	Q	S	L
	TCC	ATC	495 ACC	TGC	ACA	504 GTC	TCT		TTC	TCA			AGC	TAT	531 GGT		. CAC	540 TGG
	 S		 T		т			 G			 L			 Y		v	н	W
			549			558			567			576		•	585			594 'GGT
	V	R	Q	S	P	G	K	G	L	Ε	W	L	G	Λ	I	W	S	G
	GG.A			GAC		TAA	GCA		TIC	ATA	TCC	630 : AGA		AGC			AAG	648 GAC
	G	S		D	Y.	N	A	A				ੜ	L	S	I	S	K	D

Figure 47 cont.

AAT	TCC	657 AAG	AGC	: CAA	666 GTT			675 AAA							' AAT		
 N	 S	 К	 S	Q	 V	F	 F	K.		N	S	.T	Q	A	N	D	Т
GCC		711 TAT	TAC	TGT	720 GCC		ATG	729 GAG	AAC		738 TCG	TTT	GCT	747 TAC	TGG	GGC	756 CAA
A	I	Y	Y	C	A	R	М	Æ	N	W	S	F	A	Y	W	G	Q
GGG	ACC	765 ACG	GTC	ACC	774 GTC	TCC	GAA	, 00	CCC		792 CCT	AGC	ACC	801 CCC	CCT	GGC	810 AGC
G	T	T	v	T	V	s	E	F	P	K	P	s	T	P	P	G	s
AGT	GGT	819 GAA	CTG	GAA	828 GAG	CIG		837 AAG	CAT	CTT	846 AAA	GAA	CTT	855 CTG	 AAG	GGC	864 CCC
S	G	E	L	E	E	L	L	K	Н	L	K	E	L	L.	K	G	P
CGC	AAA	873 GGC	GAA	CTC	882 GAG	GAA	CTG	891 CTG	AAA	CAT	900 CTG	AAG	GAG	909 CTG	CTT	AAA	918 GGT
R	K	G	Ε	L	E	E	L	L	K	Н	L	K	E	L	L	K	G
GGG	AGC	927 GGA	GGC	GCG	936 CCG	GCA			TCA		954 TCT	ACA		963 AAA	ACA		972 CTA
G	S	G	G	А	P	A	P	T	S	s	S	Т	K	K	T	Q	L
CAA	CTG	981 GAG	CAT	TTA	990 CTG	CTG	GAT	999 TTA	CAG		1008 ATT	TTG		LO17 GGA	ATT		1026 AAT
Q	L	E	H	L	L	L	D	L	Q	M	I	L	N	G	I	N	N
TAC	_	L035 AAT	CCC		L044 CTC	ACC			CTC		1062 TTT	AAG		1071 TAC	ATG		LO80 AAG
Y	K	N	. Þ	K	L	T	R	M	L	T	F	K	F.	Y	М	P	K
	GCC		GAA		1098 AAA	CAT	CTT		TGT	CTA			GAA			CCT	
K	Α	T 3	E	L	K	Ĥ	L	·Q	C.	L	E	E	E	L	K	P	L
GAG		GTG			TTA												GAC
Ε	E	V	L	N	L	A	Q	S	K	N	F	H	Ļ	R	P	R	D
TTA	ATC	AGC	AAT	ATC	1206 AAC	GTA	ATA	GTT	CTG				GGA	TCT		ACA	ACA
L					N					E	L	K	G	S	Ε	Т	Т
	ATG		GAA	TAT	1260 GCT	$\mathtt{G}\!$	GAG	ACA	GCA	ACC	ATT	GTA	G4A	TTT		AAC	AGA
					A												

Figure 47 cont.

	-	1305			1314		:	1323			1332		:	1341		-	1350
TYCG	لىس⊽_	DOC	بليلية	TGT	CAA	AGC	ATC	ATC	TCA	ACA	CIG	ACT	GAC	GTC	CAT	CAC	CAT
7.7		æ	됴	C	Ω	S	I	I	S	\mathbf{T}	L	${f T}$	D	V	H	H	H
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		1359			مــــ	$\overline{}$											
CAC	CAT	CAC	TGA	TAA	GTC	GAC											
H	H	H	*	*	T												

Figure 48:

		5 ur C																
	E	CoRI	_	_														
5 '	GA	אריד A	ר ארע ארע	9 C ATG		18			. 5	7		. 3	6		45			
			- -	ATG		166	AGC	TGI	TA '	C AT	C CT	C TT	TTG	GTA	GCA	ACA	GCT	54 aca
			· .	M	G	W	S	C	·	 T	— г		 L					
			6-	,				_		_	1.,	r	با	V	A	T	A	${f T}$
	GGT	GT2	63 CAC	; TCC	CAT	72	C3 C	Oma	81	L		90)		99			100
				TCC			CAG	C1G	ACC	CAC	G TC	r car	AAA	TTC	ATG	TCC	ACA	TCA
	G	V	H	S	D	I	Q	L	\mathbf{T}	0	: S		·		~			
			117													_	_	S
	GTA	GGA	GAC	AGG	GTC	AGC	GTC	אכיכי	135			144			153			162
												AGT	CAG	AAT	GIG	GGT	ACT	AAT
	V	G	D	R	V	S	V	T	С	K	A	S	0	 N				
			171			100											T	N
	GTA	GCC	TGG	TAT	CAA	CAG	AAA	CCA	189 GGG	$C\Delta\Delta$	מטליטט	198	222		207			216
	 V	Δ	 Ta7										AAA	GCA	CTG	ATT	TAC	TCĞ
•	·	Α.	VV	Y	Q	Q	K	P	G	Q	S	P	K	A	L	I	Y	 S
			225			224												
	GCA	TCC	TAC	CGG	TAC 2	AGT (GGA (GTC	CCT	GAT	CGC	252 TTY	ACA (300	261	~~		270
	A	S	 У		 V	 -							·		AG1. (GGA :	ICT (3GG
			_	R	1	3	G	V	P	D	R	F	${f T}$	G	S	G	S	G
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				ACT (CTC A	CC A	TC A	AGC A	TAF	GTG	CAG	TCT	GAA (SAC 1	אנה כ 175	ברא כ	3	24
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-				CAA T	AT A	AC A	GC I	'AT C	CG (CTC .	ACG	TTC	GGT G	CT Ğ	GG A	.CC A	د AG C	78 TC
	F	С	Q	Q	Y]	N,	S	Y	P	 L		 E	G .		-			
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C	GAG A	ATC A	AA (GT G	ST G	96 37 C	או יוב	~ 4	05 66 6			414		42	23		4	32
-				GT G						3GC (3GC (GGC 1	CC G	GT G	GT G	GT GO	T TC	ŢŢ
	E	I	K	G (G (3 (3 . :	S (G	G	G	 G	s (-			
		4	41		40	. ^											3 5	
C	AG G	TG A	AA O	TG CZ	AG GA	G TC	'A GO	4: A C	59 יידי כי	cc c	ל היחי	168		47	77		48	16
-													AG CC	C TC	A C	AG AG	CI	$\mathcal C$
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			`	T	V	5	G	F	' 5	S]	L '	T S	S Y	G	V	Н	W	-
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ری		SC CA	G TC	T CC	A GGZ	A AAC	GG'	T CI	G GA	G TO	3G €	rg ga	A CTY	מידע בי מידע בי	י עריי כ	7 300	594	1
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		- 60	3		610													
		- AC	а GA 	C TAT	raa '	' GCA	. GCI	TIC	AT.	A TO	C AG	A CT	G AGO	YTA :	, ACC	ממי	648	:
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Fig	ure	48	con	ıt.				_									
AAT	TCC	657 AAG	AGC								684 AGT						702 ACA
N	S	K	S	Q	V	Ť.	F	K.	M	N	S	L	Q	А	N	D	T
GCC	ATA	711 TAT	TAC	TGT	720 GCC				AAC		738 TCG		GCT	747 TAC	TGG	GGC	756 CAA
A	I	Y	Y	С	А	R	M	E	N	W	S	F	A	Y	W	G	Q
GGG	ACC	765 ACG	GTC	ACC	774 GTC	TCC	GAA	783 TTC	ACC	CCG	792 CTG	GGT	GAC	801 ACC		CAC	810 ACC
G	${f T}$	Т	V	$^{-}$ T	V	S	E	F	T	P	L	G	D	$^{}$ T	 T	H	 Т
TCC	GGA	819 AAA	CCA	CTG	828 GAT	GGA	GAA	837 TAT		ACC	846 CTT		ATC	855 CGT		CGT	
S	G	K	P	L	D	G	E	Y	F	T	L	Q	I	R	G	R	E
CGC	TTC	873 GAG	ATG	TTC	882 CGA		CTG	V	GAG		900 TTG	GAA				GCC	918 CAG
R	F	E	M	F	R	E	L	N	E	Α	L	E	L	K	D	A	Q
GCT	GGG	927 AAG	GAG	CCA	936 GGG	GGG	AGC	945 GGA		GCG	954 CCG	GCA	CCT	963 ACT	TCA	AGT	972 TCT
А	G	K	E	, P	G	G	S	G	G	A	P	А	P	T	S	S	s
ACA	AAG	981 AAA	ACA	CAG	990 CTA				CAT		1008 CTG	CTG	GAT	1017 TTA	CAG	ATG	1026 ATT
T	K	K	T	Q	L	Q	L	E	Н	L	L	L	D .	r L	Q	_	I
TTG		1035 GGA	ATT			TAC			CCC		1062 CTC		AGG	1071 ATG	CTC	ACA	1080 TTT
L	N	G	I	N	N	Y	K	N	P	K	L	T	R	М	L	T	F
AAG 	TTT			CCC			GCC	ACA		CTG	1116 AAA 	CAT	CTT		TGT	CTA	
K	F	. Y ,	M	P	K	K	A	· T	E	L	K	Н	L	Q	С	L	Ε
GAA		1143 CTC	AAA 			GAG					1170 TTA	GCT	CAA	1179 AGC	AAA	AAC	1188 TTT
E	E	L	K	P	L	E	Ē	V	L	N	L	A	Q	S	K	N	F
CAC		1197 AGA	CCC		1206 GAC	TTA	ATC	1215 AGC			1224 AAC	GTA		1233 GTT			1242 CTA
Н	L	R	P	R	D	L				I	N	V	I	V	L	E	L
AAG		1251 TCT	GAA		1260 ACA				GAA		1278 GCT			1287 ACA	GCA		1296 ATT
K	G	S	E	${f T}$			М		E	Y	Α	D	Ε		А	T	I

Figure 48 cont.

	-	1305			1314		:	1323			1332			1341			1350
GTA	GAA.	TTT	CTG	AAC	AGA	TGG	ATT	ACC	TTT	TGT	CAA	AGC	ATC	ATC	TCA	ACA	CTG
V	E	F	L	N	R	W	I	\mathbf{T}	F	C	Q	S	I	I	S	\mathbf{T}	L
											- Sa	H					
	_	1359			1368			1377			تتے						
ACT	GAC	GTC	CAT	CAC	CAT	CAC	CAT	CAC	TGA	TAA	GTC	GAC					
\mathbf{T}	D	V	H	H	Н	н	Н	H	*	*							

Figure 49:

	1154	10																				
5'	Eco GAA	TTC	` A	CC Z	ATG	GGA	18 TGG	AGC	TGT	27 ATC	A:	rc c	CTC	36 TTC	TT	G G	ra c	45 3CA	ACA	GCI		54 CA
					 М	G	W	S	С	I	. :	I	L	F	L	•	V	A	\mathbf{T}	Α	7	r
	GGT	GT				GAT	72 ATC	CAG	CTG	81 ACC	C C	AG '	ICT	90 CCA	GC	A A	TC 2	99 ATG	TCT	GCF		08 CT
	-			 Н	 S		 I	Q	 L								I	M	S	Α	:	S
							126 ACC			179				144	 TC	A A	GT	153 GTT	AGT	TC		62 GT
							 T									-		Λ	 s	S		s S
	P	G		Ε	K	V	T	M	.1.							•			. •		2	16
	TAC	TT	1 G C	L71 CAC	TGG	TAC	180 CAG	CAG	AAC	TC	A C	GT		TCC	CC		AA 	CIC	TGG	AT 	Г Т 	'AT'
4	 Y	 L		н	. W	Y	Q	Q	K	S		G	A	S	3	Ρ	K	L	W	Ī		Y
	AGC	AC	'A 5	225 ICC	AAC	TTG	234 GCI	TCI	GG!	GT	C	CT	GCT	252 CG		rc i	AGT	261 GGC	AGT	- GG	. 2 G T	:70 :CT
			. <u> </u>				 A			 V		P	A	R		F	s	G	S	G	+	S
							288 CTC			20	7			30 GA	6 G G	CT (GAA	315 GAT	GCT	r GC		324 ACT
																						т
**	G	,	r	S	Y	S	L									А						
-	TAT	r T	AC.	333 TGC	CAG		TAC	AG'		T T	AC .	CCG	TAC	AC	G T				GGG			378 AAG
	 Y		 Y		Q	 Q	Y	s	G		Y	P	Y	Γ	•	F	G	G	G	,	Γ	K
							39 r GG	_		11	15			41 C GC	.4 ;C (GC	TCC	423	3			432 GGT
							 G												 G			
	L					-		_			E O			11	58			47	7 7	· C C	ÇΨ.	486 TCA
	TC	T C	AG	GTC	G AA	A CT	G CA															
	 S		0	V	K	 L			Ξ :	5 ·	G	A	. E	: :	L	V	K	P	G	÷	A	S
	•			•		C TG	.50 	14 4 G G	T T	5 CT 6	13 GC	TAC	c ac	5 C C	22 TC	ACC	AGK	53 TA	1 C TC	G I	TG	540 CAC
									 A													H
•							5 F	- 0			567			5	76			58	5			594 CCC
	ŢĆ	3G (GTG	AA	G CA 																	CCC
	V	Ň	V	K	: (2 7	v :	₽	G ,	R	G	L	·	Ξ	W	I	G	F	ξ .	<u>t</u>	D	P
	Α	ΑT	AGT	60 : GG	13 FT G0	T A	CT A	12 AG T	AC C	AT (621 GAG	AA	G T	rc P	\AG	A.G.C	. A.A.	63 G G(89 CC A	CA (CTG	648 ACT
	-						 T															
		N	5	_	<i>-</i>	_	•		-													

i	gure	49	CO	nt.																		
	GTA	GAC	6: A	57 AA :	CCC	TCC	666 AGC	ACA	GCC	67 TA	5 C A	TG ·	CAG	684 CTC	AGO	2 A		693 CTG	ACA		70 G	
	 V	D		K	P	S	S	- Т	A	Y		M	Q	L	S		S	L	T	S	I	Ξ
	GAC	TC	7 r G	11 CG	GTC	TAT	720 TAT	TGT	GCZ	72 A AC	9 A T	GG	GAC	738 TAC	TG(G G		747 CAA 	GGG	ACC	: A	
	D	S		 А	Ÿ	Y	Y	С	A	F	3	W	D	Y	W		G	Q	G	T		T
	GTC	AC	7 C G	65 TC	TCC	TCC	774 GGA	ACC	CCC	78 3 C1	33 17G (GT	GAC	792 ACC	AC	C C	AC	801 ACT	AGT	GG2		10 AA
	v	 T		V	-	s	G	T	Þ	I		G	D	${f T}$	τ		H	T	S	G		K
	CCA	CT	9 G 0	319 3AT	GGA	GAA	828 TAT	TTC	: AC	81 C C	37 FF (CAG	ATC	846 CGT	GG	ig (CGT	855 GAG		TT		64 AG
	 P	 L		D	G	 E	Y	F	T	:	L .	Q	I	R	C	}	R	E	R	F		E
	ATG	Lī	S C	373 CGA	GAG	CIG	882 AAT	GAC	G GC	8. C T	91 TG	gaa	CTC	900 AA) 5 G.F	T.	GCC	909 CAG		GG		918 AAG
	 M	 F		 R	 E	L	N	E			 L	 Е	L	K	I)	A	Q	A	G		K
	GAG	; C(: :A (927 GGG	GGG	TCC	93 <i>6</i> : GG2	5 A. GG	r Go	9 T G	45 GT	AGC	ACC	954 CA	1 A. G	rg '	TGC	963 ACC	GGC) AC		972 GAC
	 E		 -	 G	 G	 S	_ . G				G	s				Λ: 		Т	G	7		D
-	_	: A	A.G.	981 CTG	- CGC	CTC	990 C CC) r GC	C AC	ST C	99 CCC	GAG	ACC	100 CA	8 C C	TG	GAC	1017 ATC	CTY	C CC		026 CAC
	 M		 K		 R	 L	 P			 S	P	 E				 L	D	M				н
						TG	104	1		1(IG ()53 CAG	GG#	A AA	106 C CI		ΑA		1071 ACC				080 CCC
						 C				· V				 I		E	L	т				P
						C CT	100	_		1	107		TA T	111 C C	.6 4G C	AG	GTC	112 G CA	5 G GG	C T		.134 GTG
	 m	 				 L			 -	 L	- - -	 D	 I	 : (- 2	 Е		 Q		 ;	 Ү	
					_	.C AA				1	161			111	70			117 G CT	9 G CG	G A	TT	L188 GTG
						 I N																
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						2 I																
						AC A	10	60		-	269	,		12	78			128	37			1296
						AC AZ N																
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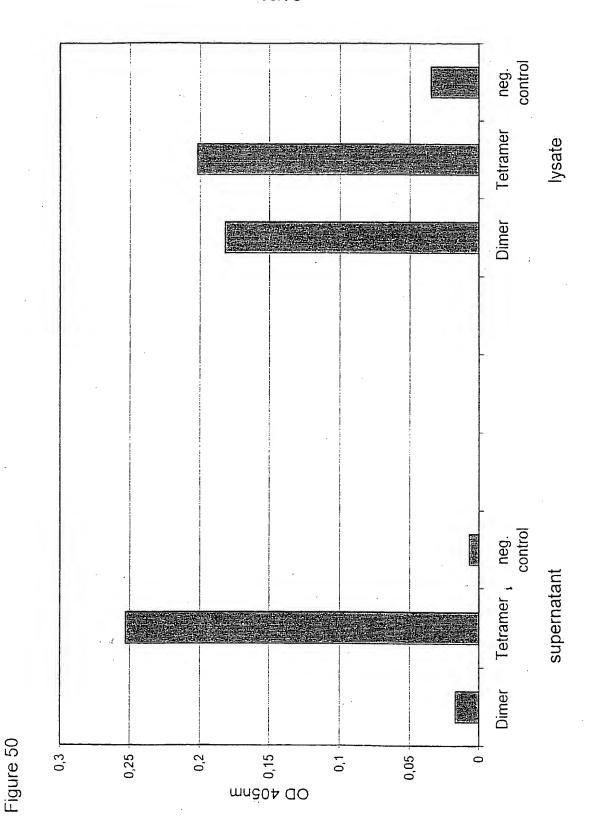
Figure 49 cont.

gure	e 49	cor	nt.				•		5									
GAG	CTG	130 CA	5 G C	TT	1 CGA	314 AGC	CIC	ACA	GAG	ATC	TTG	1332 AAA 	GGA	GGG	1341 GTC	TTG	1 ATC 	L350 CAG
E	L	Q	 !	L	R	s	L	T	E	I	L	K	G	G	V	L	I	Q.
CGG	AAC	135 : cc	9 C C	'AG	CTC	.368 TGC	TAC	CAG	1377 GAC	ACG	ATT	1386 TTG			1395 GAC	ATC		L404 CAC
R	N	P)	Q	L	С	Y	Q	D	T	I	Ļ	W	K	D	I	F	Н
AAG	AAC	141 : AA	.3 .C (CAG	CTG	GCT	CTC	ACA	1431 CTG	ATA	GAC	1440 ACC	AAC	CGC	1449 TCT			1458 TGC
K	N		1	Q	L	A	L	T	L	I	D	T	N	R	. S	R	A	С
CAC	. ccc	146 TO	57 FT 1	CT	CCG	1476 ATG	TGT	AAG	1485 GGC	TCC	CGC	1494 TGC	TGG	GGA	1503 GAG	-		1512 GAG
Н	P		2	S	P	М	С	K	G	S	R	C	M	G	E	S	S	E
			AG A	AGC	CTG	ACG	CGC	ACT		TGT	GCC		GGC	TGI	1557 GCC			1566 AAG
D	C		2	s	L	Т	R	T	V	С	Α	G	G	С	A	R	С	K
GG(G CC.	151 A C	75 IG (CCC	ACT	1584 GAC	TGC	TGC	CAT	GAG	CAC	TGI	GCI	GCC	1611 GGC	TGC		
G	P	:	L L	P	T	D	C	С	H	Ε	Q	С	A	A	G	C	. T	G
CC	C AA	16: G C	29 AC	TCT	GAC	1638 TGC	CIG	GCC	1647	CIC	CAC	1656 TTC	AAC	CAC	1665 AGT		ATC	1674 TGT
P			 H	s	D	C	L	A	С	L	Н	F	N	Н	S	G	I	С
GA:	G CI	16 G C	83 AC	TGO	: CCA	1692 . GCC	CTG	GT	1701 C ACC	TAC	AAC	1710 C AC	A GA	2 AC	1719 G TTT		TC	1728 C ATG
 E			н	C	P		L	V	т	Y				Т	F	E	S	M
CC	C AF	17 T C	737 ·	GAG	GGG	1746 CGC	; TAT	AC	1755 A TTC	5 Ç GG(C, GC	176 C AG	C TG	T GI	1773 G AC	3 r gcc	_TG	1782 T CCC
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P	_ ` _	 L	Q Q	P	E	Q		Q	V	F	E	т	L	E	E	I	Т	G
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 Y		 L	 Y		S	 А	W	P	D	S	L	P	D	L	s	V	F	Q
AA	.c c	TG	2169 CAA	GTA	ATC	2178 CGG	GGA	CGA	2187 ATT	CTG	CAC	2196 AAT	GGC	GCC	2205 TAC			2214 ACC
	 Г	 L			I	 R	G	R	I	L	H	N	G	A	Y	S	L	Т
CT	G C	2 AA	223 GGG	CTG	GGC	2232 ATC	.AGC	TGG	2241 CTG	GGG		2250 CGC	TCA	CTG	2259 AGG			2268 GGC
 L		 Q	G	L	G	I	s	W	L	G	L	R	S	L	R	E	L	G
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Figure 49 cont.

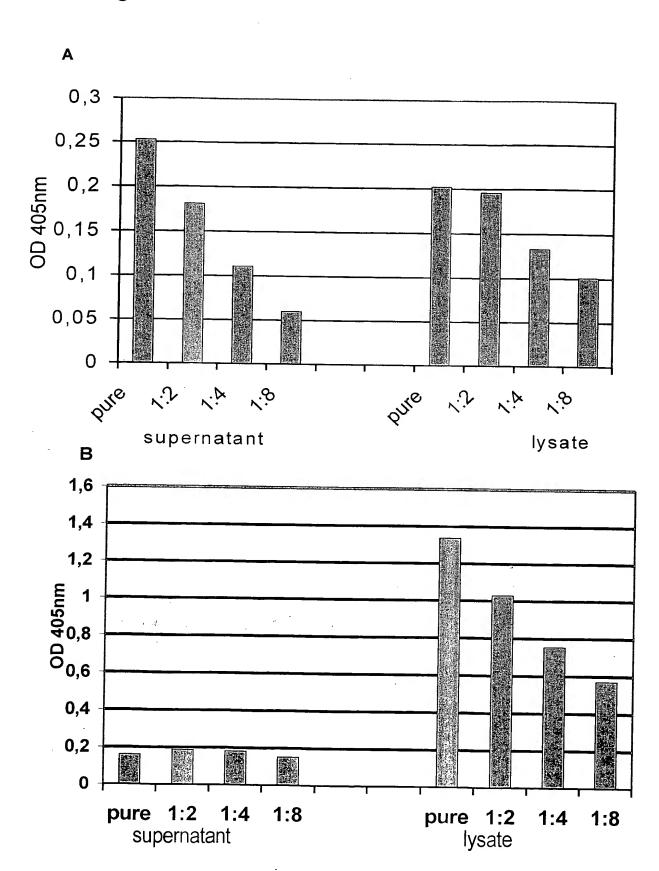
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 P	C	P CCC	ATC I	AAC N	TGC	ACC T	CAC H	TCC	TGT C	GTG V	GAC D	CTG L	GAT D	GAC D 2853	AAG K	· GGC G	TGC C 2862
 P	C	2817	I ATC	AAC N N AGA	2826	ACC T	CAC H	TCC S 2835 CTG	TGT C	GTG V	D 2844 GGG	CTG L	GAT D	GAC D 2853	AAG K	· GGC G	TGC C 2862
 P	C GCC	2817 C GAG	ATC I	AAC N	2826	ACC T	CAC H	TCC	TGT C	GTG V	GAC D	CTG L CAT	D CAT	GAC D 2853	AAG K K	GGC G CAT	TGC C 2862 CAT
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P CCC	C GCC	P 2817	I CAG	AAC N N AGA	2826	ACC T	CAC H	TCC S 2835 CTG	TGT C	GTG V	D 2844 GGG	CTG L CAT	D CAT	GAC D 2853	AAG K K	GGC G CAT	TGC C 2862 CAT
P CCC	C GCC	2817 C GAG	I CAG	AAC N N AGA	2826	ACC T	CAC H	TCC S 2835 CTG	TGT C	GTG V	D 2844 GGG	CTG L CAT	D CAT	GAC D 2853	AAG K K	GGC G CAT	TGC C 2862 CAT



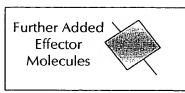
(25)

Figure51

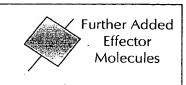


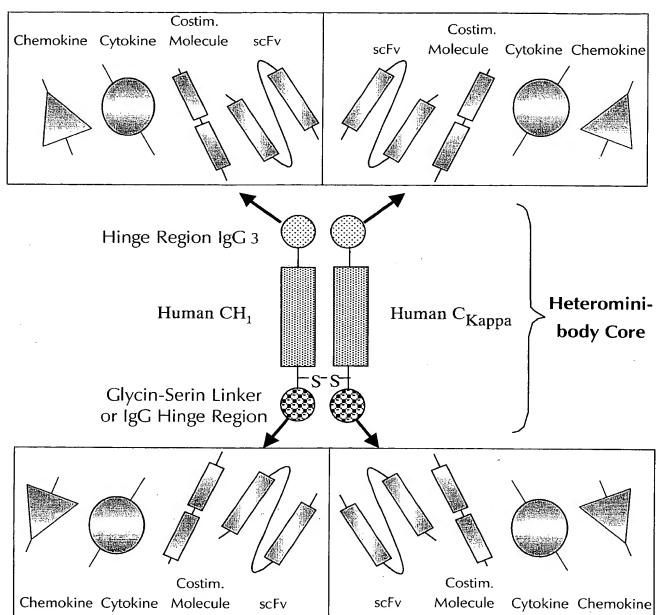


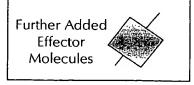
61/75



N-Terminally Linked Effector Functions







C-Terminally Linked Effector Functions

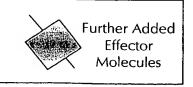


Figure 52

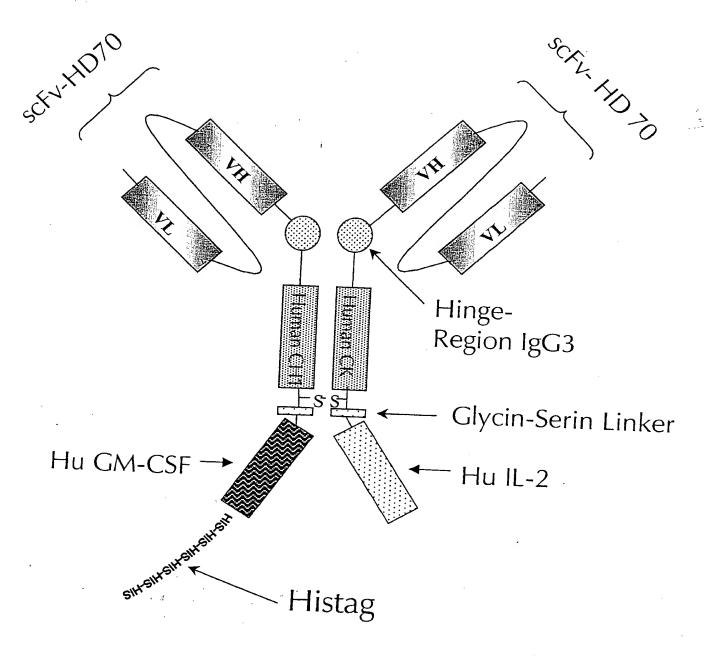


Figure 54

HD 70 scFv - CH1-GM-CSF:

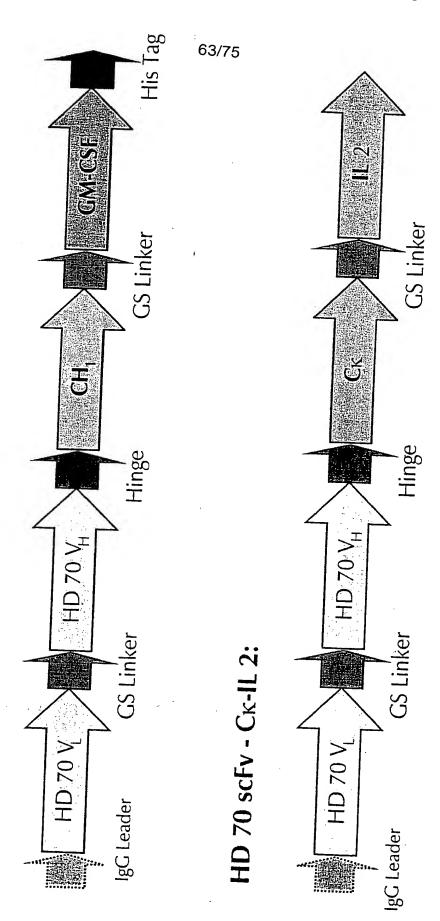


Figure 55a

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Figure 55a cont.

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901	+3 L	CCC	L CTGG GACC	CAC	P C G	S CTCC GAGG	S TCC. AGG'	AAG	S AGC TCG	T ACC1 FGG <i>F</i>	S ICTO	G G G C C	G GGC CCG	T ACA TGT	~~	A GGC: CCG:	L CCT GGA	G GGC CCCG	C TG AC	I CCT GGA	, 'GG .CC.	V TCA AGT	
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A Company of the Comp

+3 Figure 55b 67/75 M G W S EcoRI NcoI 1 TTTTTTTCTT CCATTTCAGG TGTCGTGAGG AATTCACCAT GGGATGGAGC TGTATCATCC AAAAAAAGAA GGTAAAGTCC ACAGCACTCC TTAAGTGGTA CCCTACCTCG ACATAGTAGG +3 L F L V T G V H A T A SEL ОМТ BsrGI SacI ~~~~~~ TCTTCTTGGT AGCAACAGCT ACAGGTGTAC ACTCCGAGCT CCAGATGACC CAGTCTCCAT AGAAGAACCA TCGTTGTCGA TGTCCACATG TGAGGCTCGA GGTCTACTGG GTCAGAGGTA +3 S S L S G D R V T I T CRA 121 CCTCCCTGTC TGCATCTGTA GGAGACAGAG TCACCATCAC TTGCCGGGCA AGTCAGAGCA GGAGGGACAG ACGTAGACAT CCTCTGTCTC AGTGGTAGTG AACGGCCCGT TCAGTCTCGT +3 I S S Y L N W QOK PGO PPK SwaI 181 TTAGCAGÇTA TTTAAATTGG TATCAGCAGA AACCAGGACA GCCTCCTAAG CTGCTCATTT AATCGTCGAT AAATTTAACC ATAGTCGTCT TTGGTCCTGT CGGAGGATTC GACGAGTAAA +3 Y W A S T + R - ESGVP D R F S ' G S SmaI 241 ACTGGGCATC TACCCGGGAA TCCGGGGTCC CTGACCGATT CAGCGGCAGT GAATCTGGGA TGACCCGTAG ATGGGCCCTT AGGCCCCAGG GACTGGCTAA GTCGCCGTCA CTTAGACCCT +3 T N Y T S L Q P E D A T PstI 301 CAAATTACAC TCTCACCATC AGCAGCCTGC AGCCTGAAGA TTTTGCTACT TACTTTTGTC GTTTAATGTG AGAGTGGTAG TCGTCGGACG TCGGACTTCT AAAACGATGA ATGAAAACAG +3 Q Q S D I T F G SLP QGT 361 AACAGTCTGA CAGTTTGCCG ATCACCTTCG GCCAAGGGAC ACGACTGGAC ATTCAAGGAG RLD TTGTCAGACT GTCAAACGGC TAGTGGAAGC CGGTTCCCTG TGCTGACCTG TAAGTTCCTC +3 G G G S GGG G S G G G G SE V O PvuII 421 GAGGAGGATC AGGTGGTGGT GGTAGCGGCG GCGGCGGCTC AGAGGTGCAG CTGCTCGAGT CTCCTCCTAG TCCACCACCA CCATCGCCGC CGCCGCCGAG TCTCCACGTC GACGAGCTCA +3 S G G G V V Q PGRS L R L S C A 481 CTGGGGGAGG CGTGGTCCAG CCTGGGAGGT CCCTGAGACT CTCCTGTGCA GCCTCTGGAT GACCCCCTCC GCACCAGGTC GGACCCTCCA GGGACTCTGA GAGGACACGT CGGAGACCTA +3 F T F \cdot S Y G . M H W V R Q A PGK 541 TCACCTTCAG TAGCTATGGC ATGCACTGGG TCCGCCAGGC TCCAGGCAAG GGGCTGGAGT AGTGGAAGTC ATCGATACCG TACGTGACCC AGGCGGTCCG AGGTCCGTTC CCCGACCTCA +3 W V A V I S Y DGSN $K - Y \cdot Y$ A D S V K G NdeI 601 GGGTGGCAGT TATATCATAT GATGGAAGTA ATAAATACTA TGCAGACTCC GTGAAGGGCC CCCACCGTCA ATATAGTATA CTACCTTCAT TATTTATGAT ACGTCTGAGG CACTTCCCGG S R D +3 R F T I N S K N T L Y 661 GATTCACCAT CTCCAGAGAC AATTCCAAGA ACACGCTGTA TCTGCAAATG AACAGCCTGA L Q M CTAAGTGGTA GAGGTCTCTG TTAAGGTTCT TGTGCGACAT AGACGTTTAC TTGTCGGACT

Figure 55b cont. 68/75 +3 R P Y Y M D V W G V S BspEI 781 GACCCTACTA CTACTACGGT ATGGACGTCT GGGGCCAAGG GACCACGGTC ACCGTCTCCT CTGGGATGAT GATGATGCCA TACCTGCAGA CCCCGGTTCC CTGGTGCCAG TGGCAGAGGA +3 S G T P L G D н т R T V A A P BspEI BsiWI ~~~~~ 841 CCGGAACCCC GCTGGGTGAC ACCACCCACA CCCGTACGGT GGCTGCACCA TCTGTCTTCA GGCCTTGGGG CGACCCACTG TGGTGGGTGT GGGCATGCCA CCGACGTGGT AGACAGAAGT +3 I F P P SDE QLKS G T A s v v CLL TCTTCCCGCC ATCTGATGAG CAGTTGAAAT CTGGAACTGC CTCTGTTGTG TGCCTGCTGA 901 AGAAGGGCGG TAGACTACTC GTCAACTTTA GACCTTGACG GAGACAACAC ACGGACGACT +3 N N F Y A K V Q PRE W K V D N A L O S 961 ATAACTTCTA TCCCAGAGAG GCCAAAGTAC AGTGGAAGGT GGATAACGCC CTCCAATCGG TATTGAAGAT AGGGTCTCTC CGGTTTCATG TCACCTTCCA CCTATTGCGG GAGGTTAGCC +3 G N S Q E S V T E Q D S K D S T Y S L S 1021 GTAACTCCCA GGAGAGTGTC ACAGAGCAGG ACAGCAAGGA CAGCACCTAC AGCCTCAGCA CATTGAGGGT CCTCTCACAG TGTCTCGTCC TGTCGTTCCT GTCGTGGATG TCGGAGTCGT +3 S T L T A D Y E L S K K H K CEV VYA 1081 GCACCCTGAC GCTGAGCAAA GCAGACTACG AGAAACACAA AGTCTACGCC TGCGAAGTCA CGTGGGACTG CGACTCGTTT CGTCTGATGC TCTTTGTGTT TCAGATGCGG ACGCTTCAGT +3 T H Q G L S S PVTK S F N R G E C SacI 1141 CCCATCAGGG CCTGAGCTCG CCCGTCACAA AGAGCTTCAA CAGGGGAGAG TGTTCAGGAG GGGTAGTCCC GGACTCGAGC GGGCAGTGTT TCTCGAAGTT GTCCCCTCTC ACAAGTCCTC A P T S S S T K K T Q L L E H 0 1201 GCGGTGGGTC TGCACCTACT TCAAGTTCTA CAAAGAAAAC ACAGCTACAA CTGGAGCATT CGCCACCCAG ACGTGGATGA AGTTCAAGAT GTTTCTTTTG TGTCGATGTT GACCTCGTAA LOM I L N G I N N Y K N 1261 TACTGCTGGA TTTACAGATG ATTTTGAATG GAATTAATAA TTACAAGAAT CCCAAACTCA ATGACGACCT AAATGTCTAC TAAAACTTAC CTTAATTATT AATGTTCTTA GGGTTTGAGT T F K F Y M P K K A T E L 1321 CCAGGATGCT CACATTTAAG TTTTACATGC CCAAGAAGGC CACAGAACTG AAACATCTTC GGTCCTACGA GTGTAAATTC AAAATGTACG GGTTCTTCCG GTGTCTTGAC TTTGTAGAAG +3 Q C L E · E E L K P L E E V L NLA S K XbaI 1381 AGTGTCTAGA AGAAGAACTC AAACCTCTGG AGGAAGTGCT AAATTTAGCT CAAAGCAAAA TCACAGATCT TCTTCTTGAG TTTGGAGACC TCCTTCACGA TTTAAATCGA GTTTCGTTTT +3NFHL RPR DLIS NIN VIV 1441 ACTITCACTI AAGACCCAGG GACTTAATCA GCAATATCAA CGTAATAGTI CTGGAACTAA TGAAAGTGAA TTCTGGGTCC CTGAATTAGT CGTTATAGTT GCATTATCAA GACCTTGATT +3 K G S E T T F CEY Μ Α D E TAT 1501 AGGGATCTGA AACAACATTC ATGTGTGAAT ATGCTGATGA GACAGCAACC ATTGTAGAAT TCCCTAGACT TTGTTGTAAG TACACACTTA TACGACTACT CTGTCGTTGG TAACATCTTA +3 F L N R W I T. F C Q S I I S SalI TTCTGAACAG ATGGATTACC TTTTGTCAAA GCATCATCTC AACACTAACT TGATAAGTCG AAGACTTGTC TACCTAATGG AAAACAGTTT CGTAGTAGAG TTGTGATTGA ACTATTCAGC

Figure 55b cont.

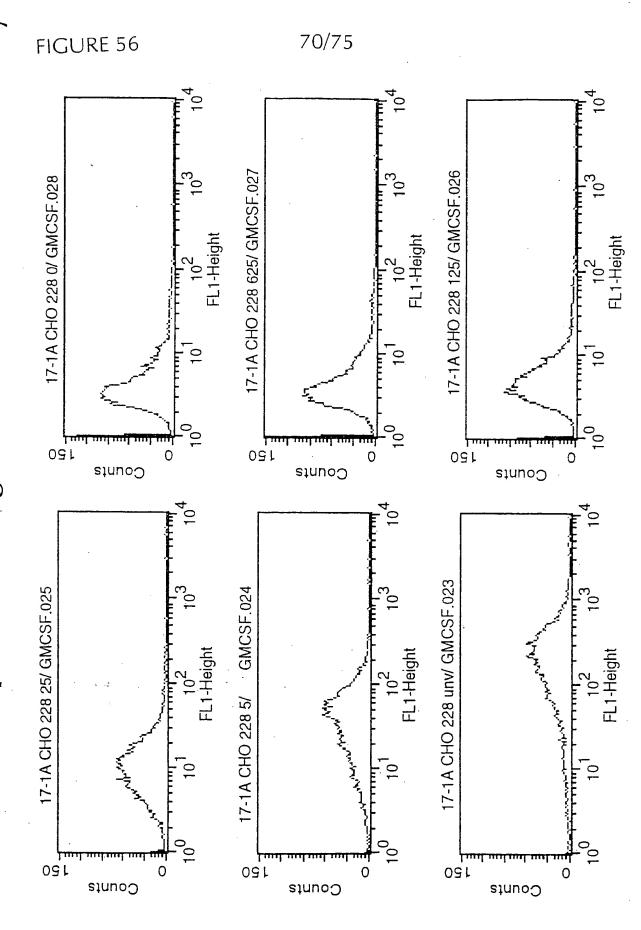
69/75

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1621 AC

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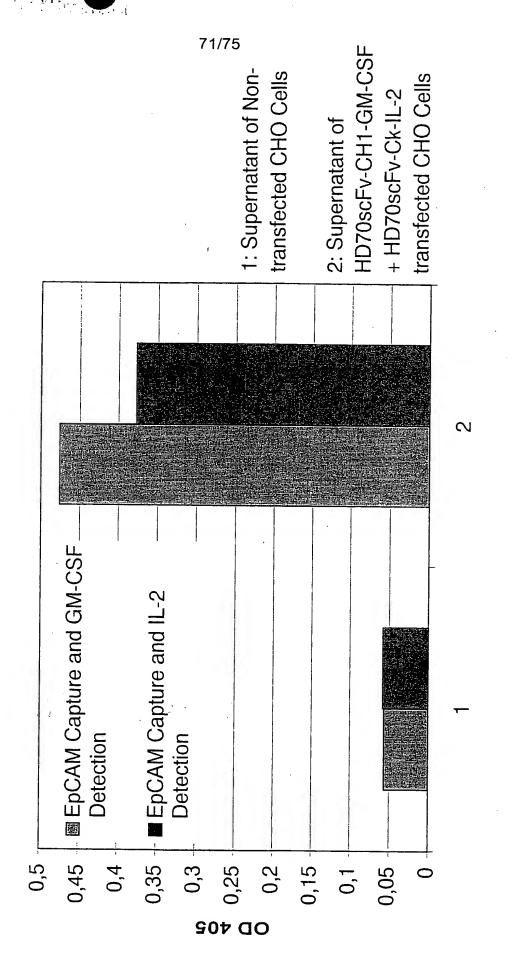
Association of EpCAM Binding and GM-CSF in Heterominibody



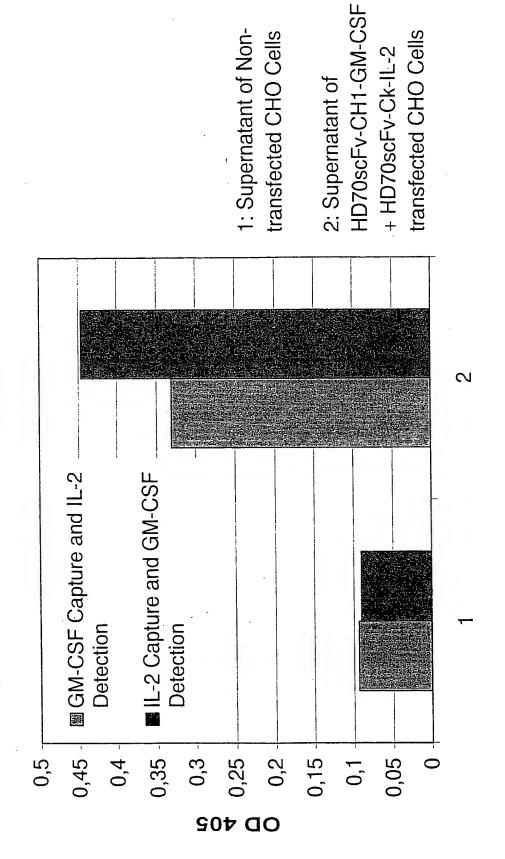
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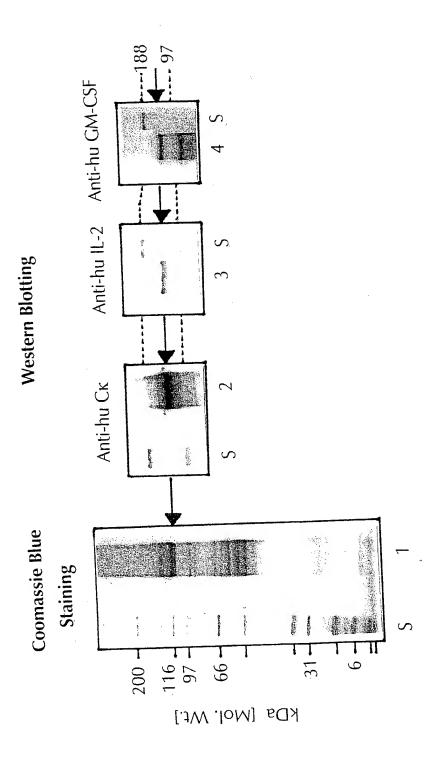
Figure 57

Physical Linkage of Anti-EpCAM Activity with IL-2 and GM-CSF



Physical Linkage of IL-2 with GM-CSF in Heterominibody





Bioactivity of GM-CSF in Heterominibody Format

74/75 0 IUGM-CSF/ml 0,3 IU GM-CSF/ml 3 IU GM-CSF/ml 300 IU GM-CSF/ml GM-CSF-IL-2-HMB 0,5 4,0 3,5 2,5 2,0 3,0 OD 420

